

1632

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/171,553A

DATE: 05/11/2001
TIME: 12:05:38

Input Set : A:\USSN 09-171,553 sequence listing.txt
Output Set: N:\CRF3\05112001\I171553A.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: GALBRAITH, DANIEL N.
4 HAWORTH, CHRISTINE
5 LEES, GILLIAN M.
6 SMITH, KENNETH T.
8 <120> TITLE OF INVENTION: PORCINE RETROVIRUS
10 <130> FILE REFERENCE: CFV-5.01
12 <140> CURRENT APPLICATION NUMBER: 09/171,553A
13 <141> CURRENT FILING DATE: 1999-02-08
15 <150> PRIOR APPLICATION NUMBER: PCT/GB97/01087
16 <151> PRIOR FILING DATE: 1997-04-18
18 <150> PRIOR APPLICATION NUMBER: GB 9702668.6
19 <151> PRIOR FILING DATE: 1997-02-10
21 <150> PRIOR APPLICATION NUMBER: GB 9608164.1
22 <151> PRIOR FILING DATE: 1996-04-19
E--> 24 <160> NUMBER OF SEQ ID NOS: 1517 (see below)
26 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORLED SEQUENCES

1557 <210> SEQ ID NO: 17
1558 <211> LENGTH: 673
1559 <212> TYPE: DNA
1560 <213> ORGANISM: Porcine retrovirus
1562 <400> SEQUENCE: 17
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1564 ataaaaagct ctaaatgccc ccgaattmca gaccctgctg gctgccagta aataggtaga 120
1565 aggtcacact tccatttggt ccaggcgctg ctatcctgac ctaagtaaga taacaggaaa 180
1566 tgagtgact aatcgcttat ctggattctg taaaactgac tggcaccata gaagaattga 240
1567 ttacacattg acagccctag tgaacctatct caactgcaat ctgtcactct gcccaggagc 300
1568 ccacgcagat gcgggacctcc ggaagctatt taaaatgatt ggtccacgga gcgcgggctc 360
1569 tcgataatttt aaaatgattg gtccatggag cgcgggctct cgaattttta aaatgattgg 420
1570 ttgtgacgc acaggccttg ttgtgaaacc cataaaagct gtcccgattc cgcaactcgg 480
1571 gcgcagctcc tctaccctcg cgtggtgtac gactgtgggc ccacgcgcg ttggaataaa 540
1572 aatcctcttg ctgtttgcat caagaccgct tctygtgagt gatttggggt gtcgcctctt 600
1573 ccgakkccgg acgaggggga ttgtctttt actgcgcttt catttgggtc gttggccggg 660
1574 aaatcctgcg acc

E--> 1579/28
E--> 1583 20/454525.1

delete at end of file

last sequence in file

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/171,553A

DATE: 05/11/2001

TIME: 12:05:40

Input Set : A:\USSN 09-171,553 sequence listing.txt
Output Set: N:\CRF3\05112001\1171553A.raw

L:1579 M:254 E: No. of Bases conflict, LENGTH:Input:28 Counted:673 SEQ:17
M:254 Repeated in SeqNo=17
L:1583 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
L:1583 M:252 E: No. of Seq. differs, <211>LENGTH:Input:673 Found:675 SEQ:17
L:24 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (15) Counted (17)